

RAW SEQUENCE LISTING ERROR REPORT



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Application Serial Number: 09/623/304
Source: 1647
Date Processed by STIC: 1/3/7/02

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FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216. PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax) PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

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Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

- 1. EFS-Bio (httm, EFS Submission User Manual ePAVE)
- 2. U.S. Postal Service: U.S. Patent and Trademark Office, Box Sequence, P.O. Box 2327, Arlington, VA 22202
- 3. Hand Carry directly to:
 - U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7th Floor, Examiner Name, Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202
 - U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 S uth Clark Place, Arlington, VA 22202
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Revised 01/29/2002



1647

Does Not Comply Corrected Diskette Needed

RAW SEQUENCE LISTING DATE:
PATENT APPLICATION: US/09/623,304 TIME:

DATE: 03/07/2002 TIME: 10:29:16

Input Set : A:\-4-1.app

Output Set: N:\CRF3\03072002\1623304.raw

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3 <110> APPLICANT: Silvia, Christopher
         Yu, Weifeng
         ICAgen, Inc.
 7 <120> TITLE OF INVENTION: Identification and Expression of Human Kir5.1
 9 <130> FILE REFERENCE: 018512-000410US
11 <140> CURRENT APPLICATION NUMBER: US 09/623,304
12 <141> CURRENT FILING DATE: 2001-02-21
14 <150> PRIOR APPLICATION NUMBER: US 60/076,612
15 <151> PRIOR FILING DATE: 1998-03-03
17 <150> PRIOR APPLICATION NUMBER: WO PCT/US99/04549
18 <151> PRIOR FILING DATE: 1999-03-02
20 <160> NUMBER OF SEQ ID NOS: 4
22 <170> SOFTWARE: PatentIn Ver. 2.1
24 <210> SEQ ID NO: 1
25 <211> LENGTH: 383
26' <212> TYPE: PRT
27 <213> ORGANISM: Homo sapiens
29 <220> FEATURE:
30 <223> OTHER INFORMATION: human Kir5.1 alpha subunit monomer of inward
        rectifier potassium channel
                                           must que location and explain - see what residue XDD represents - see
33 <220> FEATURE:
34 <221> NAME/KEY: PEPTIDE ·
35 <222> LOCATION: (351)..(383)
36 <223> OTHER INFORMATION: tail region
38 <400> SEQUENCE: 1
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41 Tyr Pro Gly Tyr Pro Pro Glu His Ile Ile Ala Glu Lys Arg Arg Ala
43 Arg Arg Arg Leu Leu His Lys Asp Gly Ser Cys Asn Val Tyr Phe Lys
                                40
45 His Ile Phe Gly Glu Trp Gly Ser Tyr Val Val Asp Ile Phe Thr Thr
47 Leu Val Asp Thr Lys Trp Arg His Met Phe Val Ile Phe Ser Leu Ser
                        70
                                             75
49 Tyr Ile Leu Ser Trp Leu Ile Phe Gly Ser Val Phe Trp Leu Ile Ala
51 Phe His His Gly Asp Leu Leu Asn Asp Pro Asp Ile Thr Pro Cys Val
               100
                                   105
53 Asp Asn Val His Ser Phe Thr Gly Ala Phe Leu Phe Ser Leu Glu Thr
                               120
                                                    125
55 Gln Thr Thr Ile Gly Tyr Gly Tyr Arg Cys Val Thr Glu Glu Cys Ser
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130

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57 Val Ala Val Leu Met Val Ile Leu Gln Ser Ile Leu Ser Cys Ile Ile
                            150
                                                155
     59 Asn Thr Phe Ile Ile Gly Ala Ala Leu Ala Lys Met Ala Thr Ala Arq
                        165
                                            170
     61 Lys Arg Ala Gln Thr Ile Arg Phe Ser Tyr Phe Ala Leu Ile Gly Met
                    180
                                        185
     63 Arg Asp Gly Lys Leu Cys Leu Met Trp Arg Ile Gly Asp Phe Arg Pro
                                    200
     65 Asn His Val Val Glu Gly Thr Val Arg Ala Gln Leu Leu Arg Tyr Thr
                                215
     67 Glu Asp Ser Glu Gly Arg Met Thr Met Ala Phe Lys Asp Leu Lys Leu
                            230
                                                235
     69 Val Asn Asp Gln Ile Ile Leu Val Thr Pro Val Thr Ile Val His Glu
                        245
                                            250
    71 Ile Asp His Glu Ser Pro Leu Tyr Ala Leu Asp Arg Lys Ala Val Ala
                    260
                                        265
     73 Lys Asp Asn Phe Glu Ile Leu Val Thr Phe Ile Tyr Thr Gly Asp Ser
                275
                                    280
W--> 75 Thr Gly Thr Ser His Gln Ser Arg Ser Ser Tyr Val Pro Arg Xaa Ile
    76
           290
                                295
                                                    300
    77 Leu Trp Gly His Arg Phe Asn Asp Val Leu Glu Val Lys Arg Lys Tyr
                            310
                                                315
                                                                     320
    79 Tyr Lys Val Asn Cys Leu Gln Phe Glu Gly Ser Val Glu Val Tyr Ala
                        325
                                            330
    81 Pro Phe Cys Ser Ala Lys Gln Leu Asp Trp Lys Asp Gln Gln Leu His
                                        345
                    340
    83 Ile Glu Lys Ala Pro Pro Val Arg Glu Ser Cys Thr Ser Asp Thr Lys
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    85 Ala Arg Arg Arg Ser Phe Ser Ala Val Ala Ile Val Ser Ser Trp
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    90 <211> LENGTH: 1509
    91 <212> TYPE: DNA
    92 <213> ORGANISM: Homo sapiens
    94 <220> FEATURE:
    95 <223> OTHER INFORMATION: human Kir5.1 alpha subunit monomer of inward
              rectifier potassium channel
    98 <220> FEATURE:
    99 <221> NAME/KEY: unsure
    100 <222> LOCATION: (1279)
    101 <223> OTHER INFORMATION: n = a, g, c or t
    103 <400> SEQUENCE: 2
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    105 agcagctatc atattatcaa tgcggacgca aaatacccag gctacccgcc agagcacatt 120
    106 atagctgaga agagaagagc aagaagacga ttacttcaca aagatgqcag ctqtaatgtc 180
    107 tacttcaage acatttttgg agaatgggga agetatgtgg ttgacatett caccactett 240
    108 gtggacacca agtggcgcca tatgtttgtg atattttctt tatcttatat tctctcgtgg 300
    109 ttgatatttg gctctgtctt ttggctcata gcctttcatc atggcgatct attaaatgat 360
    110 ccagacatca caccttgtgt tgacaacgtc cattetttca caggggcctt tttgttctcc 420
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PATENT APPLICATION: US/09/623,304 TIME: 10:29:16

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111 ctagagaccc aaaccaccat aggatatggt tatcgctgtg ttactgaaga atgttctgtg 480 112 gccgtgctca tggtgatcct ccagtccatc ttaagttgca tcataaatac ctttatcatt 540 113 ggagctgcct tggccaaaat ggcaactgct cgaaagagag cccaaaccat tcgtttcagc 600 114 tactttgcac ttataggtat gagagatggg aagctttgcc tcatgtggcg cattggtgat 660 115 tttcggccaa accacgtggt agaaggaaca gttagagccc aacttctccg ctatacagaa 720 116 gacagtgaag ggaggatgac gatggcattt aaagacctca aattagtcaa cgaccaaatc 780 117 atcctggtca ccccggtaac tattgtccat gaaattgacc atgagagccc tctgtatgcc 840 118 cttgaccgca aagcagtagc caaagataac tttgagattt tggtgacatt tatctatact 900 119 ggtgattcca ctggaacatc tcaccaatct agaagctcct atgttccccg araaattctc 960 120 tggggccata ggtttaatga tgtcttggaa gttaagagga agtattacaa agtgaactgc 1020 121 ttacagtttg aaggaagtgt ggaagtatat gccccctttt gcagtgccaa gcaattggac 1080 122 tggaaagacc agcagctcca catagaaaaa gcaccaccag ttcgagaatc ctgcacgtcg 1140 123 gacaccaagg cgagacgaag gtcatttagt gcagttgcca ttgtcagcag ctggtgaaaa 1200 124 ccctgaggag accaccactt tcgccacaca tgaatatagg gaaacacctt atcagaaagc 1260 W--> 125 tetecetgae tttaaacang aateetetgt wgaateecaa atgttagtee taaaattgea 1320 126 attatgaggg ctaccactga atcattttat ctttcagcca atcaagtcgt tgtaaacgtg 1380 127 gcttttttga aagtgttatg gctatgtttt atgatgatgc tgggtaagta gagtaagtta 1440 128 aacttggtaa aagataatct aaaaattcca tagttctcag ttattaaaat ttttcttgtt 1500 129 ccggaattc 131 <210> SEQ ID NO: 3 132 <211> LENGTH: 24 133 <212> TYPE: DNA 134 <213> ORGANISM: Artificial Sequence 136 <220> FEATURE: 137 <223> OTHER INFORMATION: Description of Artificial Sequence:primer 139 <400> SEQUENCE: 3 140 cctaaqqqca caqcaaaqaa tqaq 24 142 <210> SEQ ID NO: 4 143 <211> LENGTH: 20 144 <212> TYPE: DNA 145 <213> ORGANISM: Artificial Sequence 147 <220> FEATURE: 148 <223> OTHER INFORMATION: Description of Artificial Sequence:primer 150 <400> SEQUENCE: 4 151 gtgtggcgaa agtggtggtc 20

RAW SEQUENCE LISTING ERROR SUMMARY PATENT APPLICATION: US/09/623,304

DATE: 03/07/2002 TIME: 10:29:18

Input Set : A:\-4-1.app

Output Set: N:\CRF3\03072002\1623304.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the $\langle 220 \rangle$ to $\langle 223 \rangle$ fields of each sequence which presents at least one n or Xaa.

Seq#:1; Xaa Pos. 303
Seq#:2; N Pos. 1279

VERIFICATION SUMMARY

DATE: 03/07/2002

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Input Set : A:\-4-1.app
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L:75 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:288 L:125 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:1260